

Serial Number: 09/591737

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: ~~The above~~ corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/591,737

DATE: 06/21/2000
TIME: 11:05:33

Input Set: I591737.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

1 <110> APPLICANT: Curiel et al.
2 <120> TITLE OF INVENTION: Immunomodulation by genetic modification of
3 dendritic cells and B cells
4 <130> FILE REFERENCE: D6167CIP
5 <140> CURRENT APPLICATION NUMBER: US/09/591,737
6 <141> CURRENT FILING DATE: 2000-06-12
7 <150> EARLIER APPLICATION NUMBER: 60/102,257
8 <151> EARLIER FILING DATE: 1998-09-29
9 <160> NUMBER OF SEQ ID NOS: 1
10 <210> SEQ ID NO 1
11 <211> LENGTH: 9
12 <212> TYPE: PRT
13 <213> ORGANISM: Unknown
14 <220> FEATURE:
15 <221> NAME/KEY: DOMAIN
16 <222> LOCATION:
17 <223> OTHER INFORMATION: RGD motif that binds specifically to _v_3 and
18 _v_5 integrins
19 <400> SEQUENCE: 1
20 Cys Asp Cys Arg Gly Asp Cys Phe Cys
21 5

PAGE: 2

VERIFICATION SUMMARY
PATENT APPLICATION US/09/591,737

DATE: 06/21/2000
TIME: 11:05:33

Input Set: I591737.RAW

Line ? Error/Warning

Original Text

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/591,737

DATE: 06/21/2000
TIME: 11:05:33

Input Set: I591737.RAW

PREVIOUSLY ERRORED SEQUENCES-EDITED

```
1  <210> 1
2  <211> 9
3  <212> PRT
4  <213> Unknown
5  <220>
6  <221> DOMAIN
7  <222>
8  <223> RGD motif that binds specifically to _v_3 and
9      _v_5 integrins
10 <400> 1
11      Cys Asp Cys Arg Gly Asp Cys Phe Cys
12                      5
```

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/591,737DATE: 06/16/2000
TIME: 19:51:19

Input Set: I591737.RAW

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

1 <110> Curiel et al.
2 <120> Immunomodulation by genetic modification of
3 dendritic cells and B cells
4 <130> D6167CIP
5 <140> US/09/591,737
6 <141> 2000-06-12
7 <150> 60/102,257
8 <151> 1998-09-29
9 <160> 1

Does Not Contain
Corrected Sequence

ERRORED SEQUENCES FOLLOW

10 <210> 1
E--> 11 <211> 9
12 <212> PRT
13 <213> Unknown
14 <220>
15 <221> DOMAIN
16 <222>
17 <223> RGD motif that binds specifically to _v_3 and
18 _v_5 integrins
19 <400> 1
20 Cys Asp Cys Arg Gly Asp Cys Phe Cys
21 5
E--> 22

SEQ

Input Set: I591737.RAW

Line	? Error/Warning	Original Text
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11	E Input 9, Calc Seq.Length 10 differ	
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		<211> 9
--	--	---------

22	E Wrong Amino Acid Designator	
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		SEQ 1
--	--	-------